locisels leigt



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98 64 64	197 124 124 124	296 185 185 185
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	101	201 + + + + + + + 250 + + + + + + 300
NIM MOUSE I-k ^l RAT I-kBα PIG I-kBα	NIMI MOUSE I-kBa RAT I-kBa PIG I-kBa	NIMI MOUSE I-KBa RAT I-KBa PIG I-KBa

FIGURE 1A

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

394 250 250 250	494 267 267 267	593 314 314 314
400 KGRLC	500 HQSRL 	600 HRRR L L
+ TPEQCKHSI	+ APFRILEB IQQ	+ - STOGEKESINRKIJSHRR -GGORITIL GGORIITIL
+ ATMAVECHIN	+ + + + + + + + + + + + + + + + + + +	+ TSSTSKSTQ
	+ + + + + + + + + + + + + + + + + + +	550
301 + + + + + + + +	+ #KGICEFIVI	+ + + + + +
350	450 AQAAMETAEN	550 - QRYMEIQEITA ESYDIESE ESYDIESE ESYDIESE
AMRKEPQL) AMRKEPQL) ADLQUEDLA ADLQUEDLA ADLQUEDLA ADLQUEDLA	AQRLFPTE2	
+ 	+ LDLENKVAL	+ :: : LACGEDOTA LPESEDE LPESEDE LPESEDE
+ .: \\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	+ AADELKWITL	+ :- : .umencentro .umencom .umencom
+ :. : XYD_LK_DI_ADN XYD_LK_LDI_ADN XYD_LG_ADN XYD_LG_ADN XYD_LG_ADN XYD_LG_ADN XYD_LG_ADN XYD_LG_ADN XYD_LG_ADN	401 + + + + + + + + VEILEQEDKREQIPRDVPPSFAVAADELKWITL	+ + + + + + + + + + + + + + + + + + +
+ : .VAYCNVKTZ STHGYLAT STHGYLGT STHGYLGT STHGYLGT	+)HIDKREQIP	+
301 : CALEEP C.LEEP C.LEEP	•	
301	NIM MOUSE I-kBa RAT I-kBa PIG I-kBa	NIMI MOUSE I-kBa RAT I-kBa PIG I-kBa

FIGURE

ICOISEIS LEIGH

	Rice-1	:	33	+ + +ALD+ DIELVKL++ + +LDDA A+H+AV +CN IRRMRRALDAADIELVKLMVMGEGLDLDDALAVHYAVQHCN	155
	NIM1	:	327	PRGYTVLHVAAMRKEPQLILSLLEKGASASEATLEGRT 364 P G T LH+AA P ++ LL+ A + T +G T	l
	Rice-1	:	215	PTGKTALHLAAEMVSPDMVSVLLDHHADXNFRTXDGVT 328	3
	NIM1	:	267	VSNVHKALDSDDIELVKLLLKEDHTNLDDACALHFAVAYCN + + +ALD+ DIELVKL++ + +LDDA A+H+AV +CN	307
	Rice-2	:	33	IRRMRRALDAADIELVKLMVMGEGLDLDDALAVHYAVQHCN	155
				RNPRGYTVLHVAAMRKEPQLILSLLEK 351 R P T LH+AA P ++ LL++	
	Rice-2	:	208	RRPDSKTALHLAAEMVSPDMVSVLLDQ 288	
				VSNVHKALDSDDIELVKLLLKEDHTNLDDACALHFAVAYCN + + +ALD+ DIELVKL++ + +LDDA A+H+AV +CN	
e Fé	Rice-3	:	33	IRRMRRALDAADIELVKLMVMGEGLDLDDALAVHYAVQHCN	155
fij	NIM1	:	325	RNPRGYTVLHVAAMRKEPQLILSLLEK 351 R P T LH+AA P ++ LL++	
	Rice-3	:	208	RRPDSKTALHLAAEMVSPDMVSVLLDQ 288	
[-2	NIM1	:	267	VSNVHKALDSDDIELVKLLLKEDHTNLDDACALHFAVAYCN + + +ALD+ DIELVKL++ + +LDDA A+H+AV +CN	307
	Rice-4	:	33	IRRMRRALDAADIELVKLMVMGEGLDLDDALAVHYAVQHCN	155
	NIM1	:	327	PRGYTVLHVAAMRKEPQLI 345 P G T LH+AA P ++	
	D 3 4		21 5	DMGIZMAT III AA DIGIZADDIGI 071	

Rice-4: 215 PTGKTALHLAAEMVSPDMV 271

NIM1 : 267 VSNVHKALDSDDIELVKLLLKEDHTNLDDACALHFAVAYCN 307

FIGURE 2